281278USOPCT.ST25 SEQUENCE LISTING

Soto-Jara, Claudio Maundrell, Kinsey <110> USE OF PRION CONVERSION MODULATING AGENTS <120> 281278US0PCT <130> 10/560,978 <140> 2005-12-16 <141> PCT/EP04/51170 <150> <151> 2004-06-18 <150> EP 03101795.7 <151> 2003-06-19 <160> <170> PatentIn version 3.3 <210> 4563 <211> <212> PRT <213> Homo sapiens <400> Met Asp Pro Pro Arg Pro Ala Leu Leu Ala Leu Leu Ala Leu Pro Ala 1 10 15 Leu Leu Leu Leu Leu Ala Gly Ala Arg Ala Glu Glu Met Leu 20 25 30 Glu Asn Val Ser Leu Val Cys Pro Lys Asp Ala Thr Arg Phe Lys His
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Ile Ser Ala Leu Leu Val Pro Pro Glu Thr Glu Glu Ala Lys Gln Val 165 170 175 Leu Phe Leu Asp Thr Val Tyr Gly Asn Cys Ser Thr His Phe Thr Val 180 185 190 Lys Thr Arg Lys Gly Asn Val Ala Thr Glu Ile Ser Thr Glu Arg Asp 195 200 205 Leu Gly Gln Cys Asp Arg Phe Lys Pro Ile Arg Thr Gly Ile Ser Pro 210 220 Leu Ala Leu Ile Lys Gly Met Thr Arg Pro Leu Ser Thr Leu Ile Ser 225 230 235 240 Ser Ser Gln Ser Cys Gln Tyr Thr Leu Asp Ala Lys Arg Lys His Val 245 250 255 Ala Glu Ala Ile Cys Lys Glu Gln His Leu Phe Leu Pro Phe Ser Tyr 260 265 270 Asn Asn Lys Tyr Gly Met Val Ala Gln Val Thr Gln Thr Leu Lys Leu 275 280 285 Glu Asp Thr Pro Lys Ile Asm Ser Arg Phe Phe Gly Glu Gly Thr Lys 290 295 300 Lys Met Gly Leu Ala Phe Glu Ser Thr Lys Ser Thr Ser Pro Pro Lys 305 310 315 320 Gln Ala Glu Ala Val Leu Lys Thr Leu Gln Glu Leu Lys Lys Leu Thr 325 330 335 Ile Ser Glu Gln Asn Ile Gln Arg Ala Asn Leu Phe Asn Lys Leu Val $340 \hspace{1.5cm} 345 \hspace{1.5cm} 350$ Thr Glu Leu Arg Gly Leu Ser Asp Glu Ala Val Thr Ser Leu Leu Pro 355 360 365 Gln Leu Ile Glu Val Ser Ser Pro Ile Thr Leu Gln Ala Leu Val Gln 370 380 Cys Gly Gln Pro Gln Cys Ser Thr His Ile Leu Gln Trp Leu Lys Arg 385 390 395 400 Val His Ala Asn Pro Leu Leu Ile Asp Val Val Thr Tyr Leu Val Ala 405 410 415 Leu Ile Pro Glu Pro Ser Ala Gln Gln Leu Arg Glu Ile Phe Asn Met 420 425 430

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Tyr Asn Thr Lys Asp Lys Ile Gly Val Glu Leu Thr Gly Arg Thr 2000 2010 Leu Ala Asp Leu Thr Leu Leu Asp Ser Pro Ile Lys Val Pro Leu 2015 2020 2025 Leu Leu Ser Glu Pro Ile Asn Ile Ile Asp Ala Leu Glu Met Arg 2030 2040 Asp Ala Val Glu Lys Pro Gln Glu Phe Thr Ile Val Ala Phe Val 2045 2055 Lys Tyr Asp Lys Asn Gln Asp Val His Ser Ile Asn Leu Pro Phe 2060 2070 Phe Glu Thr Leu Gln Glu Tyr Phe Glu Arg Asn Arg Gln Thr Ile 2075 2080 2085 Ile Val Val Glu Asn Val Gln Arg Asn Leu Lys His Ile Asn 2090 2095 2100 Ile Asp Gln Phe Val Arg Lys Tyr Arg Ala Ala Leu Gly Lys Leu 2105 2115 Pro Gln Gln Ala Asn Asp Tyr Leu Asn Ser Phe Asn Trp Glu Arg 2120 2130 Gln Val Ser His Ala Lys Glu Lys Leu Thr Ala Leu Thr Lys Lys 2135 2140 2145 Tyr Arg Ile Thr Glu Asn Asp Ile Gln Ile Ala Leu Asp Asp Ala 2150 2160 Lys Ile Asn Phe Asn Glu Lys Leu Ser Gln Leu Gln Thr Tyr Met 2165 2170 2175 Ile Gln Phe Asp Gln Tyr Ile Lys Asp Ser Tyr Asp Leu His Asp 2180 2185 2190 Leu Lys Ile Ala Ile Ala Asn Ile Ile Asp Glu Ile Ile Glu Lys 2195 2200 2205 Leu Lys Ser Leu Asp Glu His Tyr His Ile Arg Val Asn Leu Val 2210 2215 2220 Lys Thr Ile His Asp Leu His Leu Phe Ile Glu Asn Ile Asp Phe 2225 2230 2235 Asn Lys Ser Gly Ser Ser Thr Ala Ser Trp Ile Gln Asn Val Asp 2240 2245 2250

Thr Lys Tyr Gln Ile Arg Ile Gln Ile Gln Glu Lys Leu Gln Gln 2255 2260 2265 Leu Lys Arg His Ile Gln Asn Ile Asp Ile Gln His Leu Ala Gly 2270 2280 Lys Leu Lys Gln His Ile Glu Ala Ile Asp Val Arg Val Leu Leu 2285 2290 2295 Asp Gln Leu Gly Thr Thr Ile Ser Phe Glu Arg Ile Asn Asp Val 2300 2305 2310 Leu Glu His Val Lys His Phe Val Ile Asn Leu Ile Gly Asp Phe 2315 2320 2325 Glu Val Ala Glu Lys Ile Asn Ala Phe Arg Ala Lys Val His Glu 2330 2340 Leu Ile Glu Arg Tyr Glu Val Asp Gln Gln Ile Gln Val Leu Met 2345 2350 2355 Asp Lys Leu Val Glu Leu Thr His Gln Tyr Lys Leu Lys Glu Thr 2360 2365 2370 Ile Gln Lys Leu Ser Asn Val Leu Gln Gln Val Lys Ile Lys Asp 2375 2380 2385 Tyr Phe Glu Lys Leu Val Gly Phe Ile Asp Asp Ala Val Lys Lys 2390 2395 2400 Leu Asn Glu Leu Ser Phe Lys Thr Phe Ile Glu Asp Val Asn Lys 2405 2416 Phe Leu Asp Met Leu Ile Lys Lys Leu Lys Ser Phe Asp Tyr His 2420 2430 Gln Phe Val Asp Glu Thr Asn Asp Lys Ile Arg Glu Val Thr Gln 2435 2440 2445 Arg Leu Asn Gly Glu Ile Gln Ala Leu Glu Leu Pro Gln Lys Ala 2450 2455 2460 Glu Ala Leu Lys Leu Phe Leu Glu Glu Thr Lys Ala Thr Val Ala 2465 2470 2475 Val Tyr Leu Glu Ser Leu Gln Asp Thr Lys Ile Thr Leu Ile Ile 2480 2485 2490 Asn Trp Leu Gln Glu Ala Leu Ser Ser Ala Ser Leu Ala His Met 2495 2500 2505

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Leu Met Asp Glu Thr Met Lys Glu Leu Lys Ala Tyr Lys Ser Glu Leu 85 90 95

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Gly Gln Ser Thr Glu Glu Leu Arg Val Arg Leu Ala Ser His Leu Arg 145 150 155 160

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